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## Product datasheet for TP301251M

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## PSMC5 (NM_002805) Human Recombinant Protein

## Product data:

Product Type:
Description:

Species:
Expression Host:
Expression cDNA Clone
or AA Sequence:

Recombinant Proteins
Recombinant protein of human proteasome (prosome, macropain) 26S subunit, ATPase, 5 (PSMC5), $100 \mu \mathrm{~g}$

Human
HEK293T
>RC201251 protein sequence
Red=Cloning site Green=Tags(s)
MALDGPEQMELEEGKAGSGLRQYYLSKIEELQLIVNDKSQNLRRLQAQRNELNAKVRLLREELQLLQEQG SYVGEVVRAMDKKKVLVKVHPEGKFVVDVDKNIDINDVTPNCRVALRNDSYTLHKILPNKVDPLVSLMMV EKVPDSTYEMIGGLDKQIKEIKEVIELPVKHPELFEALGIAQPKGVLLYGPPGTGKTLLARAVAHHTDCT FIRVSGSELVQKFIGEGARMVRELFVMAREHAPSIIFMDEIDSIGSSRLEGGSGGDSEVQRTMLELLNQL DGFEATKNIKVIMATNRIDILDSALLRPGRIDRKIEFPPPNEEARLDILKIHSRKMNLTRGINLRKIAEL MPGASGAEVKGVCTEAGMYALRERRVHVTQEDFEMAVAKVMQKDSEKNMSIKKLWK

## TRTRPLEQKLISEEDLAANDILDYKDDDDKV

## Tag: C-Myc/DDK

Predicted MW: $\quad 45.4 \mathrm{kDa}$
Concentration: $\quad>0.05 \mu \mathrm{~g} / \mu \mathrm{L}$ as determined by microplate BCA method
Purity:
$>80 \%$ as determined by SDS-PAGE and Coomassie blue staining

Recombinant protein was captured through anti-DDK affinity column followed by

Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.

25 mM Tris-HCl, 100 mM glycine, $\mathrm{pH} 7.3,10 \%$ glycerol conventional chromatography steps.
Note: $\quad$ For testing in cell culture applications, please filter before use. Note that you may experience
$\begin{array}{ll}\text { Note: } & \text { For testing in cell culture applications, please filter } \\ & \text { some loss of protein during the filtration process. }\end{array}$
Storage: $\quad$ Store at $-80^{\circ} \mathrm{C}$.
Stability:

RefSeq:
Buffer:
Preparation:

NP 002796

| Locus ID: | S705 |
| :--- | :--- |
| UniProt ID: | P62195, AOA140VJS3 |
| RefSeq Size: | 1372 |
| Cytogenetics: | $17 q 23.3$ |
| RefSeq ORF: | 1218 |
| Synonyms: | p45; p45/SUG; RPT6; S8; SUG-1; SUG1; TBP10; TRIP1 |
| Summary: | The 26S proteasome is a multicatalytic proteinase complex with a highly ordered structure <br> composed of 2 complexes, a 20S core and a 19S regulator. The 20S core is composed of 4 <br> rings of 28 non-identical subunits; 2 rings are composed of 7 alpha subunits and 2 rings are <br> composed of 7 beta subunits. The 19S regulator is composed of a base, which contains 6 |
|  | ATPase subunits and 2 non-ATPase subunits, and a lid, which contains up to 10 non-ATPase <br> subunits. Proteasomes are distributed throughout eukaryotic cells at a high concentration <br> and cleave peptides in an ATP/ubiquitin-dependent process in a non-lysosomal pathway. An <br> essential function of a modified proteasome, the immunoproteasome, is the processing of <br> class I MHC peptides. This gene encodes one of the ATPase subunits, a member of the triple-A <br> family of ATPases which have a chaperone-like activity. In addition to participation in <br> proteasome functions, this subunit may participate in transcriptional regulation since it has <br> been shown to interact with the thyroid hormone receptor and retinoid X receptor-alpha. Two |
| transcript variants encoding different isoforms have been found for this gene. [provided by |  |

## Product images:



Coomassie blue staining of purified PSMC5 protein (Cat\# [TP301251]). The protein was produced from HEK293T cells transfected with PSMC5 cDNA clone (Cat\# [RC201251]) using MegaTran 2.0 (Cat\# [TT210002]).

